C E C C C C	hanged a file from non-ASCII to ASCII hanged the margins in cases where the sequence text was "wrapped" down to the next line. dited a format error in the Current Application Data section, specifically: dited the Current Application Data section with the actual current number. The number inputterchance of the prior application data; or other dided the mandatory heading and subheadings for "Current Application Data". dited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer manged the spelling of a mandatory field (the headings or subheadings), specifically:
E CI CI	dited a format error in the Current Application Data section, specifically: dited the Current Application Data section with the actual current number. The number inputterchically oplicant was the prior application data; or other dided the mandatory heading and subheadings for "Current Application Data". dited the "Number of Sequences" (ield. The applicant spelled out a number instead of using an integer
En Ci	dited the Current Application Data section with the actual current number. The number inputterchycen oplicant was the prior application data; or other dided the mandatory heading and subheadings for "Current Application Data". dited the "Number of Sequences" (field. The applicant spelled out a number instead of using an integen
Ar Er Cl	oplicant was the prior application data; or other Idded the mandatory heading and subheadings for "Current Application Data". Idited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer
CI Ci	dited the "Number of Sequences" (ield. The applicant spelled out a number instead of using an intege
Ci	
Co	nanged the spelling of a mandatory field (the headings or subheadings), specifically:
_	
Ins	orrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	serted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	procted subheading placement. All responses must be on the same line as each subheading. If the plicant placed a response below the subheading, this was moved to its appropriate place.
In	serted colons after headings/subheadings. Headings edited included:
D	eleted extra, invalid, headings used by an applicant, specifically:
D	eleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end o page numbers throughout text; other invalid text, such as
lr	serted mandatory headings, specifically: <u>L2207-leg2</u>
С	orrected an obvious error in the response, specifically:
E	dited identifiers where upper case is used but lower case is required, or vice versa.
С	orrected an error in the Number of Sequences field, specifically:
A	"Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
De du	leted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (end to a Patentin bug). Sequences corrected:
0	ther: Jeg 2 - conected and and hos
•	

DATE: 06/27/2000

TIME: 21:14:19

Input Set : A:\Pto.amc Output Set: N:\CRF3\06272000\I424487.raw 3 <110> APPLICANT: CHOO, Yen KLUG, Aaron ISALAN, Mark 7 <120> TITLE OF INVENTION: Nucleic Acid Binding Proteins 9 <130> FILE REFERENCE: 71278/264975 11 <140> CURRENT APPLICATION NUMBER: US 09/424,487 C--> 12 <141> CURRENT FILING DATE: 2000-02-29 14 <150> PRIOR APPLICATION NUMBER: GB 9710809.6 15 <151> PRIOR FILING DATE: 1997-05-23 17 <150> PRIOR APPLICATION NUMBER: PCT/GB98/01512 18 <151> PRIOR FILING DATE: 1998-05-26 20 <160> NUMBER OF SEQ ID NOS: 2 22 <170> SOFTWARE: PatentIn Ver. 2.1 24 <210> SEQ ID NO: 1 25 <211> LENGTH: 264 26 <212> TYPE: DNA 27 <213> ORGANISM: Artificial Sequence 29 <220> FEATURE: 30 <221> NAME/KEY: CDS 31 <222> LOCATION: (1)..(264) 33 <220> FEATURE: 34 <223> OTHER INFORMATION: Description of Artificial Sequence: encoding 35 <223> OTHER INFORMATION: nucleic acid binding proteins 37 <400> SEQUENCE: 1 38 gca gaa gag aag cot ttt cag tgt cga atc tgc atg cgt aac ttc agc 39 Ala Glu Glu Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser 10 42 gat cgt act act ctt acc cgc cac acg agg acc cac aca ggc gag aag 43 Asp Arg Thr Thr Leu Thr Arg His Thr Arg Thr His Thr Gly Glu Lys 25 46 cct ttt cag tgt cga atc tgc atg cgt aac ttc agc agg agc gat aac 47 Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Arg Ser Asp Asn 40 48 35 50 ctt acg aga cac cta agg acc cac aca ggc gag aag cct ttt cag tgt 192 51 Leu Thr Arg His Leu Arg Thr His Thr Gly Glu Lys Pro Phe Gln Cys 55 52 50 54 cga atc tgc atg cgt aac ttc agg caa gct gat cat ctt caa gag cac 240 55 Arg Ile Cys Met Arg Asn Phe Arg Gln Ala Asp His Leu Gln Glu His 75 56 65 70 264 58 cta aag acc cac aca ggc gag aag 59 Leu Lys Thr His Thr Gly Glu Lys 85 63 <210> SEQ ID NO: 2 64 <211> LENGTH: 88 65 <212> TYPE: PRT 66 <213> ORGANISM: Artificial Sequence W--> 67 <220> FEATURE:

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/424,487

RAW SEQUENCE LISTING

DATE: 06/27/2000

PATENT APPLICATION: US/09/424,487

TIME: 21:14:19

Input Set : A:\Pto.amc

Output Set: N:\CRF3\06272000\I424487.raw

68 <223> OTHER INFORMATION: Description of Artificial Sequence: encoding
69 <223> OTHER INFORMATION: nucleic acid binding proteins
71 <400> SEQUENCE: 2
72 Ala Glu Glu Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser
73 1 5 10 15
75 Asp Arg Thr Thr Leu Thr Arg His Thr Arg Thr His Thr Gly Glu Lys
76 20 25 30
78 Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Arg Ser Asp Asn
79 35 40 45
81 Leu Thr Arg His Leu Arg Thr His Thr Gly Glu Lys Pro Phe Gln Cys
82 50 55 60
84 Arg Ile Cys Met Arg Asn Phe Arg Gln Ala Asp His Leu Gln Glu His
85 65 70 75 80
87 Leu Lys Thr His Thr Gly Glu Lys
88

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/424,487

DATE: 06/27/2000 TIME: 21:14:20

Input Set : A:\Pto.amc
Output Set: N:\CRF3\06272000\I424487.raw

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:67 M:283 W: Missing Blank Line separator, <220> field identifier